Sequence Listing

(1) GENERAL INFORMATION:

    (i) APPLICANT:
        (A) NAME: Max-Planck-Gesellschaft zur Foerderung der
              Wissenschaften e.V
        (B) STREET: none
        (C) CITY: Berlin
        (E) COUNTRY: Germany
        (F) POSTAL CODE: none

    (ii) TITLE OF INVENTION: Process for increasing the yield in plants

    (iii) NUMBER OF SEQUENCES: 1

    (iv) COMPUTER READABLE FORM:
        (A) MEDIUM TYPE: Floppy disk
        (B) COMPUTER: IBM PC compatible
        (C) OPERATING SYSTEM: PC-DOS/MS-DOS
        (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPA)

(2) INFORMATION FOR SEQ ID NO: 1:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 1138 base pairs
        (B) TYPE: nucleotide
        (C) STRANDEDNESS: double
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: Genomic DNA

    (iii) HYPOTHETICAL: NO

    (iv) ANTISENSE: NO

    (vi) ORIGINAL SOURCE:
        (A) ORGANISM: Agrobacterium rhizogenes

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

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AATTCGATAC GAAAAAGGCA AGTGCCAGGG CCATTTAAAA TACGGCGTCG GAAACTGGCG      60

CCAATCAGAC ACAGTCTCTG GTCGGGAAAG CCAGAGGTAG TTTGGCAACA ATCACATCAA     120

GATCGATGCG CAAGACACGG GAGGCCTTAA AATCTGGATC AAGCGAAAAT ACTGCATGCG     180

TGATCGTTCA TGGGTTCATA GTACTGGGTT TGCTTTTTCT TGTCGTGTTG TTTGGCCTTA     240

GCGAAAGGAT GTCAAAAAAG GATGCCCATA ATTGGGAGGA GTGGGGTAAA GCTTAAAGTT     300

GGCCCGCTAT TGGATTCGC GAAAGCGGCA TTGGCAAACG TGAAGATTGC TGCATTCAAG     360

ATACTTTTTC TATTTCTGG TTAAGATGTA AAGTATTGCC ACAATCATAT TAATTACTAA     420

CATTGTATAT GTAATATAGT GCGGAAATTA TCTATGCCAA AATGATGTAT TAATAATAGC     480
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AATAATAATA TGTGTTAATC TTTTTCAATC GGGAATACGT TTAAGCGATT ATCGTGTTGA      540

ATAAATTATT CCAAAAGGAA ATACATGGTT TTGGAGAACC TGCTATAGAT ATATGCCAAA      600

TTTACACTAG TTTAGTGGGT GCAAAACTAT TATCTCTGTT TCTGAGTTTA ATAAAAAATA      660

AATAAGCAGG GCGAATAGCA GTTAGCCTAA GAAGGAATGG TGGCCATGTA CGTGCTTTTA      720

AGAGACCCTA TAATAAATTG CCAGCTGTGT TGCTTTGGTG CCGACAGGCC TAACGTGGGG      780

TTTAGCTTGA CAAAGTAGCG CCTTTCCGCA GCATAAATAA AGGTAGGCGG GTGCGTCCCA      840

TTATTAAAGG AAAAAGCAAA AGCTGAGATT CCATAGACCA CAAACCACCA TTATTGGAGG      900

ACAGAACCTA TTCCCTCACG TGGGTCGCTA GCTTTAAACC TAATAAGTAA AAACAATTAA      960

AAGCAGGCAG GTGTCCCTTC TATATTCGCA CAACGAGGCG ACGTGGAGCA TCGACAGCCG     1020

CATCCATTAA TTAATAAATT TGTGGACCTA TACCTAACTC AAATATTTTT ATTATTTGCT     1080

CCAATACGCT AAGAGCTCTG GATTATAAAT AGTTTGGATG CTTCGAGTTA TGGGGTAC      1138
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